

Figure 1

CgLyse 1 MVIMEIFITGLLLGASLLLSIGPONVLVIKOGIKREGLIAVLLVCI. TSIV 50
EcYgga 1-MILPLGPQNAFVMNOGIRRQYHIMIALLCAISDL 34

CgLyse 51 FLFIAGTLGVVDLLSNAAPIVLDIMRWGGIAYLLWFAVMAAKDAMTNKVEA 100
EcYgga 35 VLICAGIFGGSALLMQSPWLLALVTWGGVAFLWYGFCAFKTAMSSNIE. 83

CgLyse 101 PQIEETEPTVPDDTPLGGSAVATDTRNRVRVEVSVDKQRVWVKPMILMAI 150
EcYgga 84LASAEVMKQGKWK.....IIATMLAV 104

CgLyse 151 VLTWLNPNAYLD~~A~~FVFIGGVGAQYGDTGRW~~I~~FAAGAFAASLIWFPLVGFG 200
EcYgga 105 ..TWLNPHVYLD~~I~~FVVLGSLGGOLDVEPKRW~~F~~ALGTISASFLWFFGLALL 152

CgLyse 201 AAALSRPLSSPKVWRWINVVAVVMTALAIKLMLMG. 236
EcYgga 153 AAWLAPRLRTAKAQRI~~I~~NLVVGCVMW~~F~~IALQ~~L~~ANDGIAHQALFS 197

Figure 2

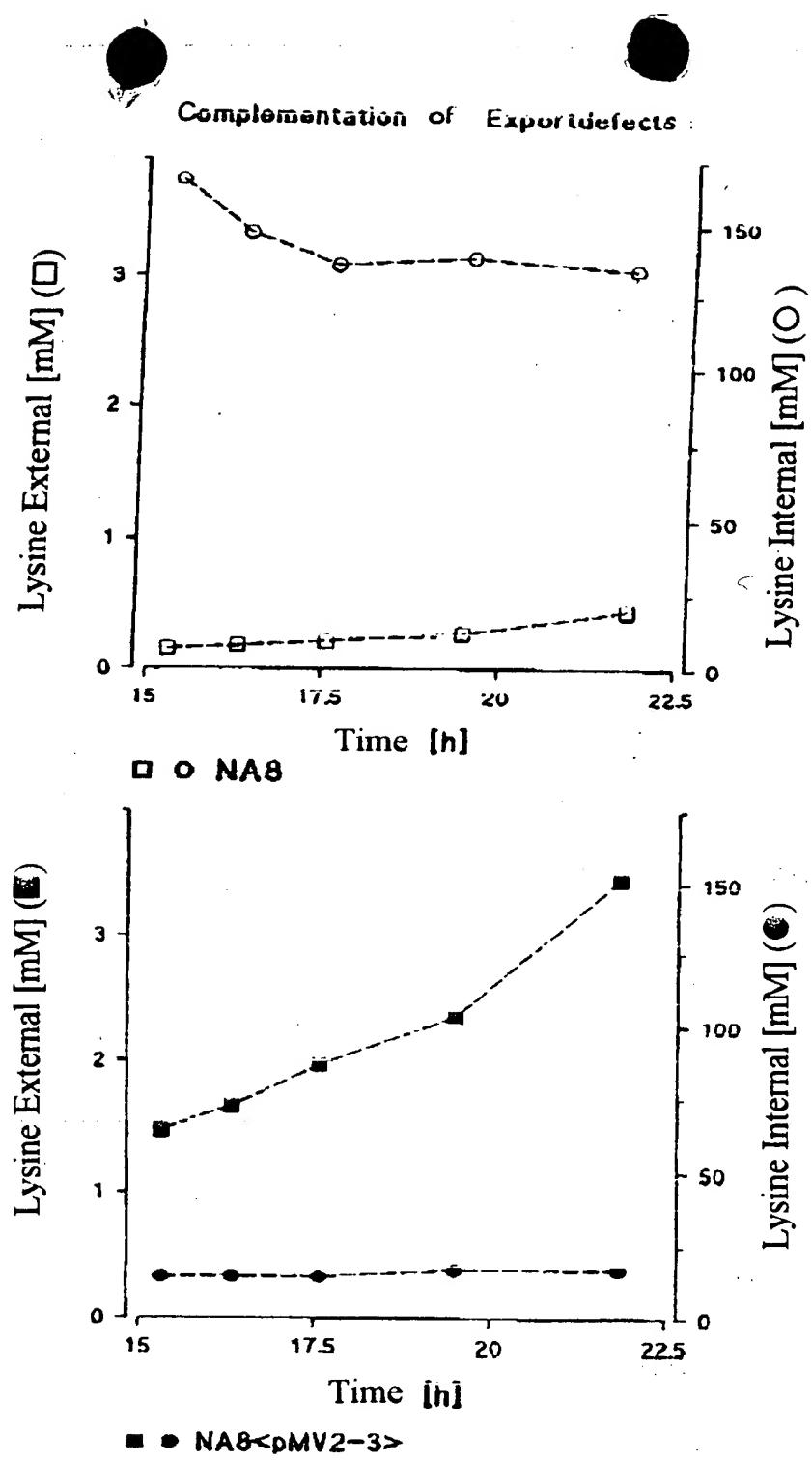


Figure 3

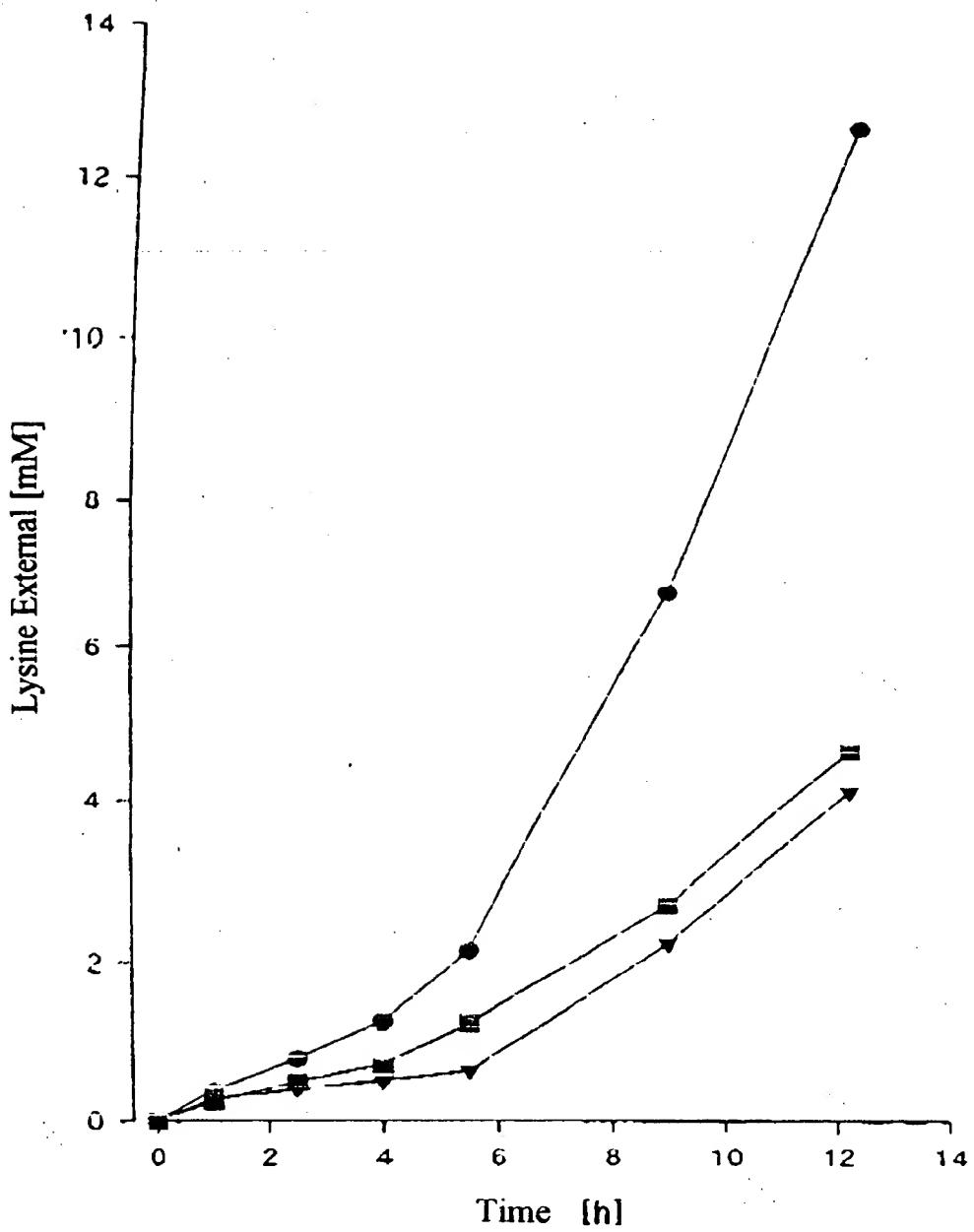


Figure 4